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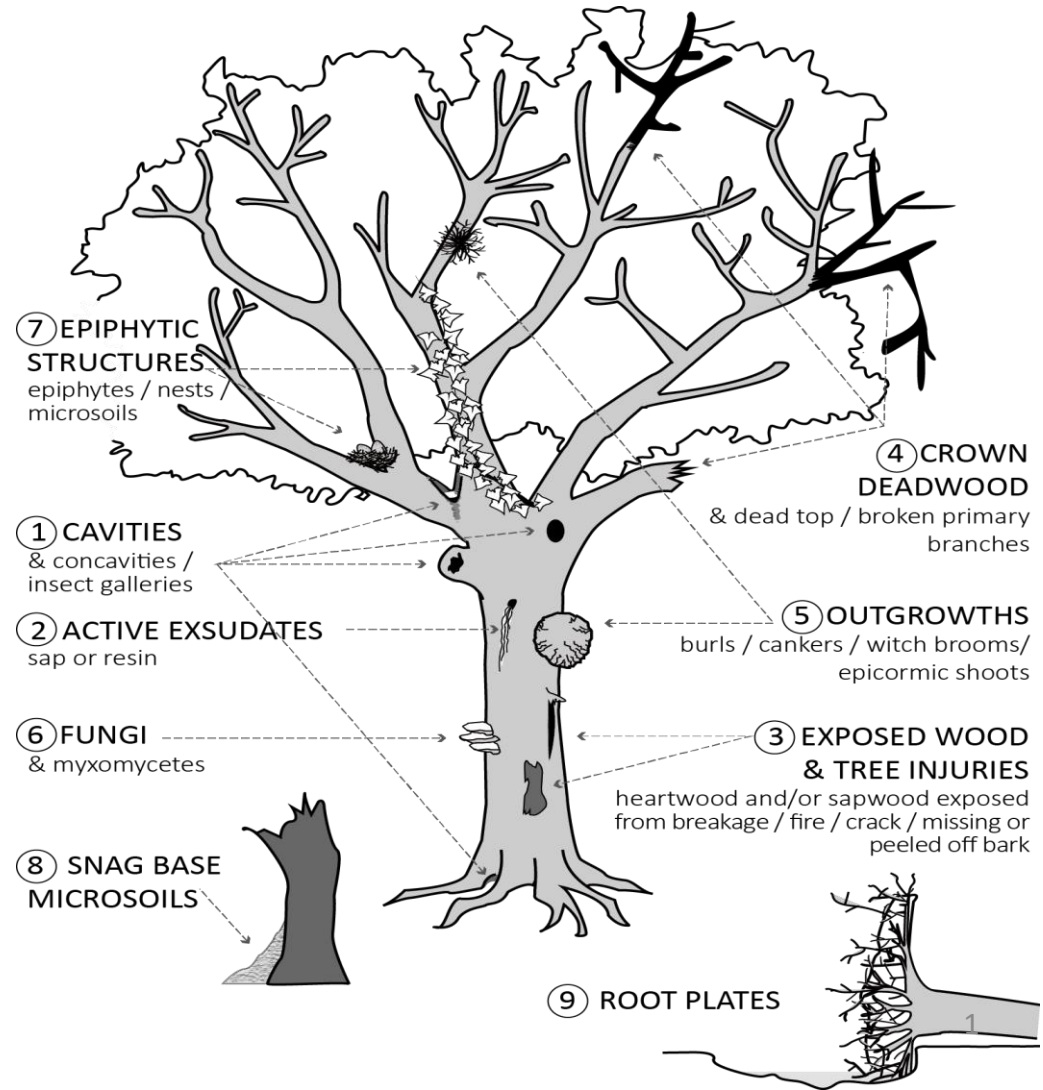
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# Tree-related microhabitats (TreMs) as key elements for forest biodiversity

Laurent LARRIEU<sup>1,2</sup>  
Christophe BOUGET<sup>3</sup>

1-INRA Toulouse    
2-CNPF-CRPFOc   
3-IRSTEA EFNO 



# TReMs are morphological singularities borne by trees; they host a wide range of taxa

- Borne by above ground parts of a tree, dead or living: base, trunk and crown
- Small to mid-sized ( $\text{cm}/\text{cm}^3 \rightarrow \text{m}/\text{m}^3$ )
- Encompassing decaying wood (=saproxylic TreMs) or not (=epixylic TreMs)
- Hosting preferentially-associated taxa

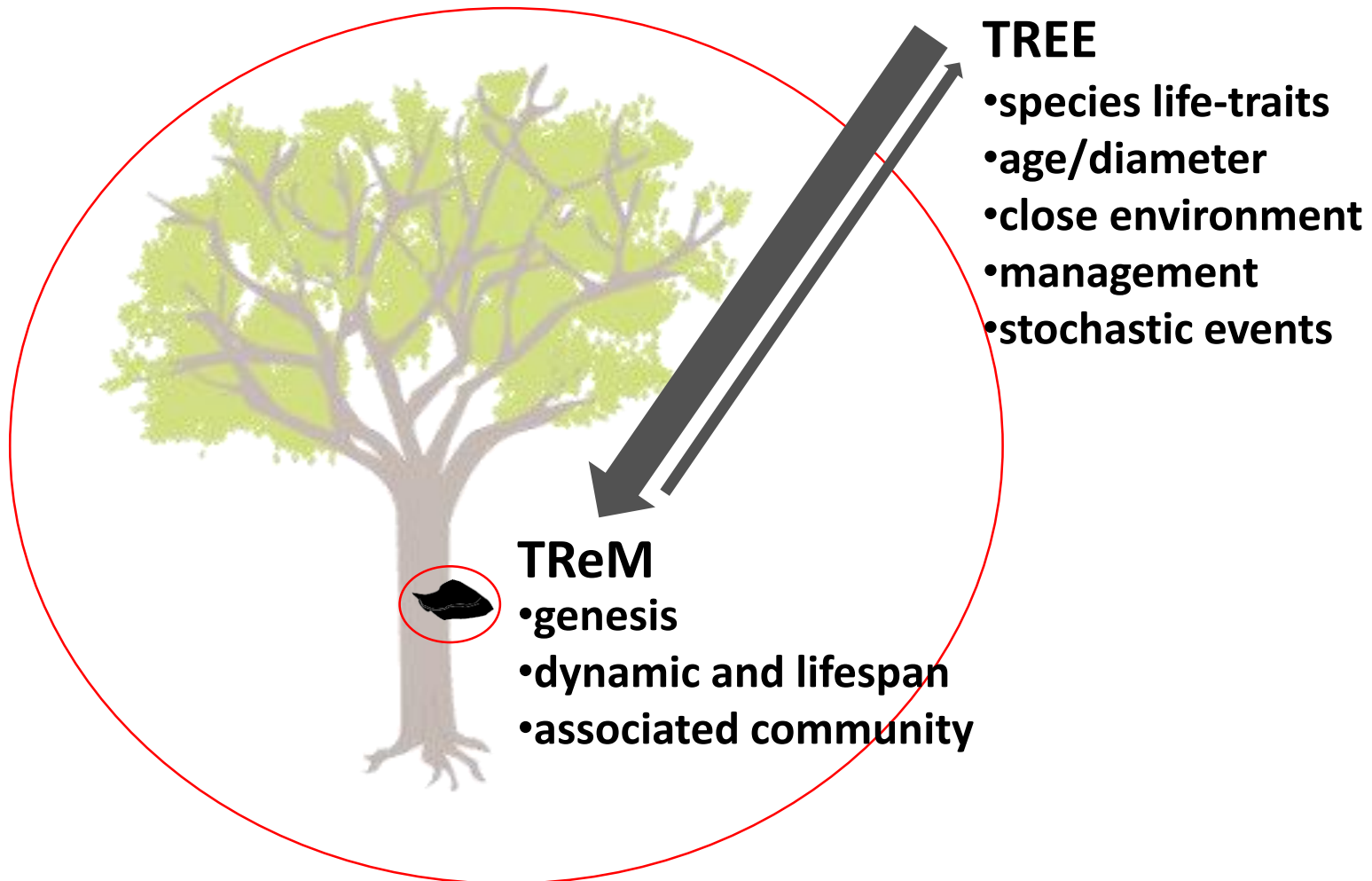


# I-TreMs as ecological items





# TreMs depend on tree characteristics



And tree vitality and life-span sometimes depends on the TreM it bears...

# TReMs are « ephemeral resource patches » (Finn 2001) »



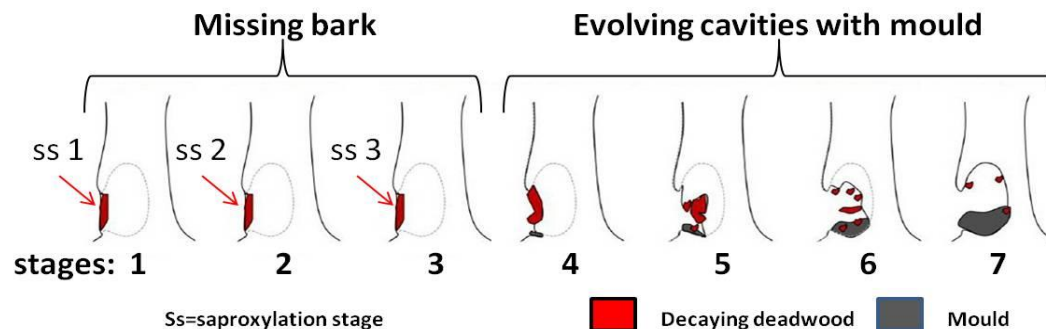
- high quality site → Dependence gradient
- spatially limited → Small size (limited by the tree size)
- temporary → TReM type “X”

**Disappearance**  
(tree removed)

**Development/change**

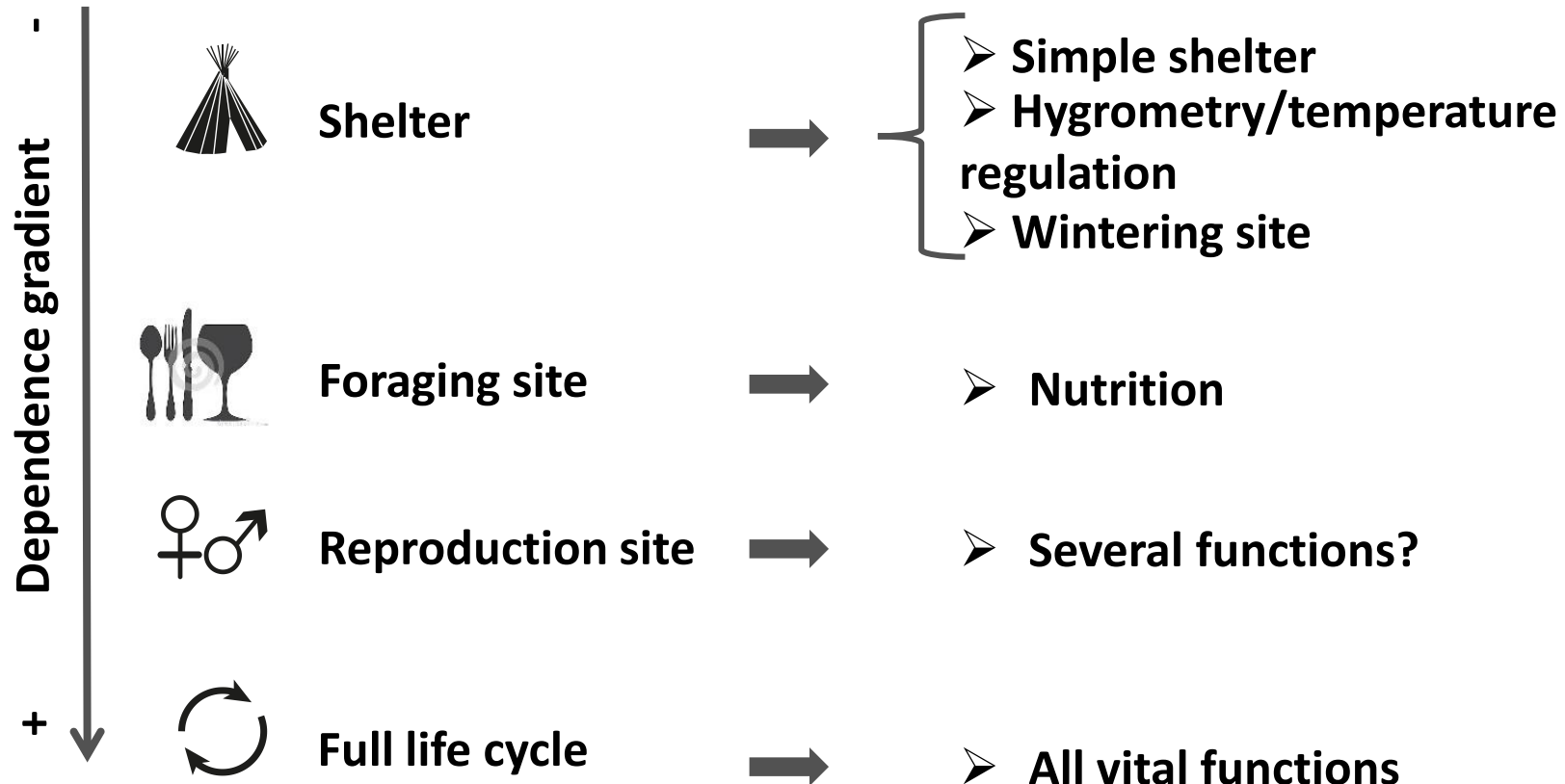
**Unavailability**  
(=“useful” period)

□ Type “X” → Type “Y”  
(=lifespan)

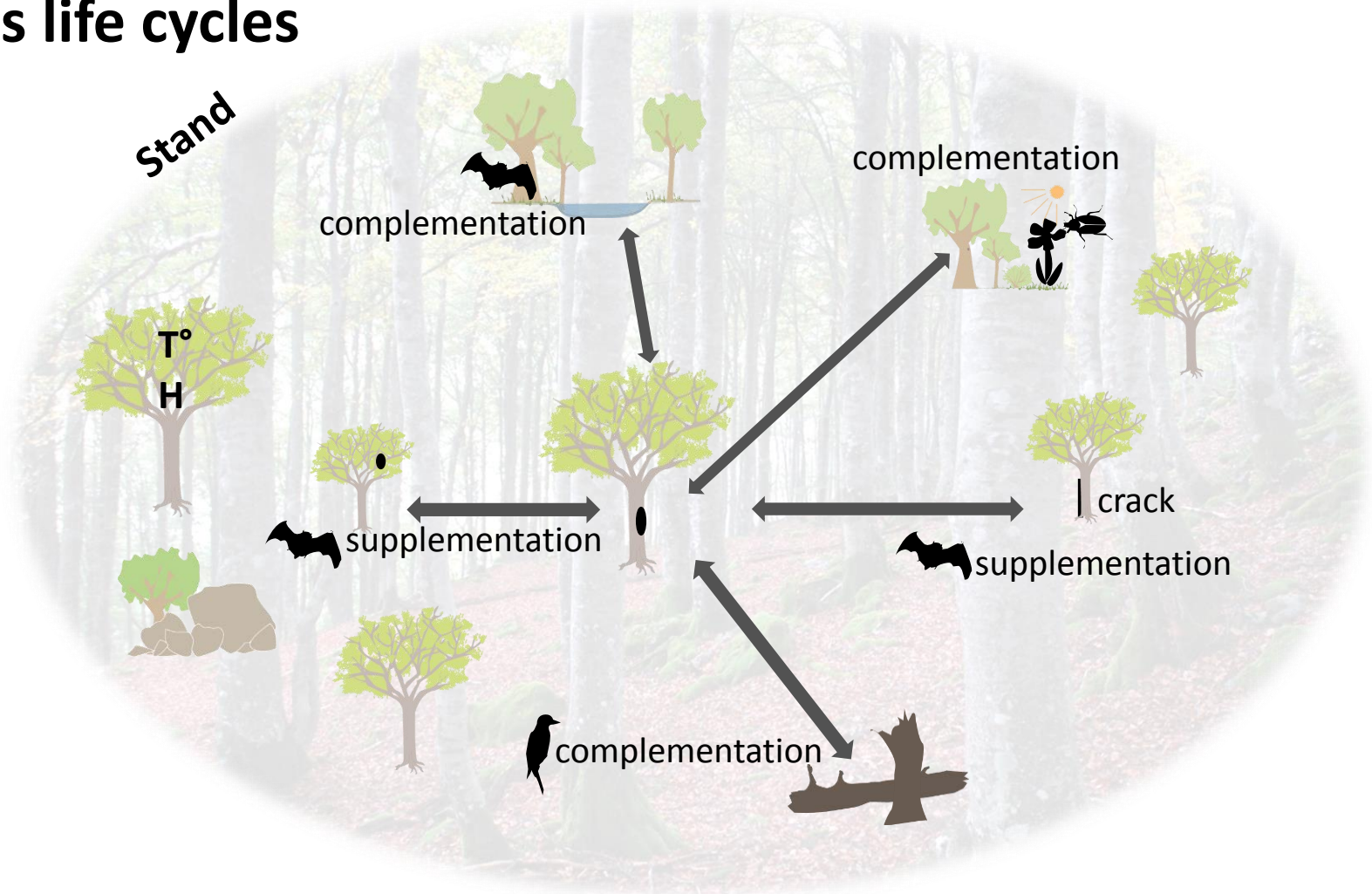





□ Living tree → Dead tree

# TreMs play a wide range of pivotal biological roles



# TReMs participate in a functional habitat network in species life cycles



Examples of complementation/supplementation resources for woodpeckers (  ), bats (  ), saproxylic beetles<sup>7</sup> (  )



# TReMs are key elements for complexity of forest ecosystems

## Dimensions of complexity (from Cadenasso et al. 2006)

### Structural heterogeneity

- ☐ Broad scale
  - Trees
  - Eco-units
  - Sylvigenetic phases
  - Sylvigenetic cycle

- ☐ Fine scale
  - **TreM density**
  - **TreM diversity**
  - **Substrate diversity within a TreM**

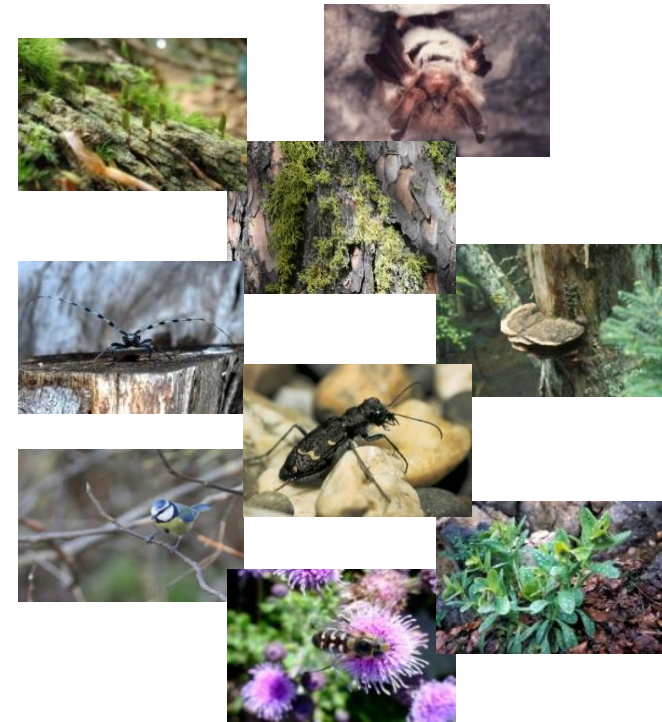
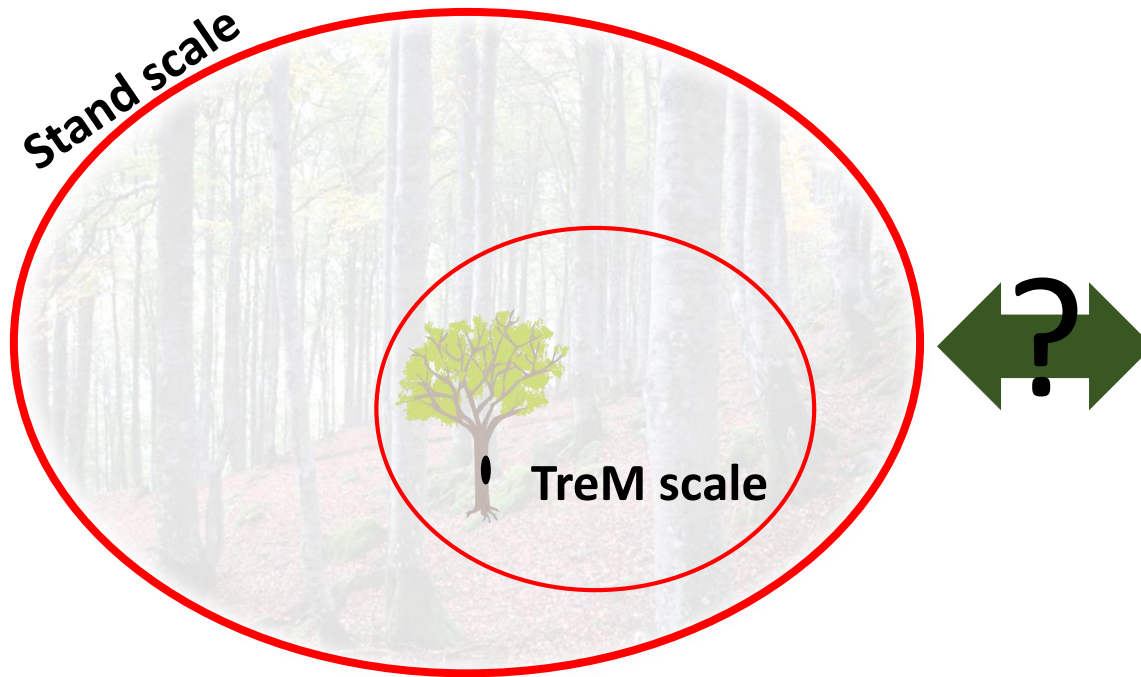
### Spatial connectivity

- **TreM spatial distribution**
- **TreMs as complementation/supplementation resources**
- **Changes in TreM profile**

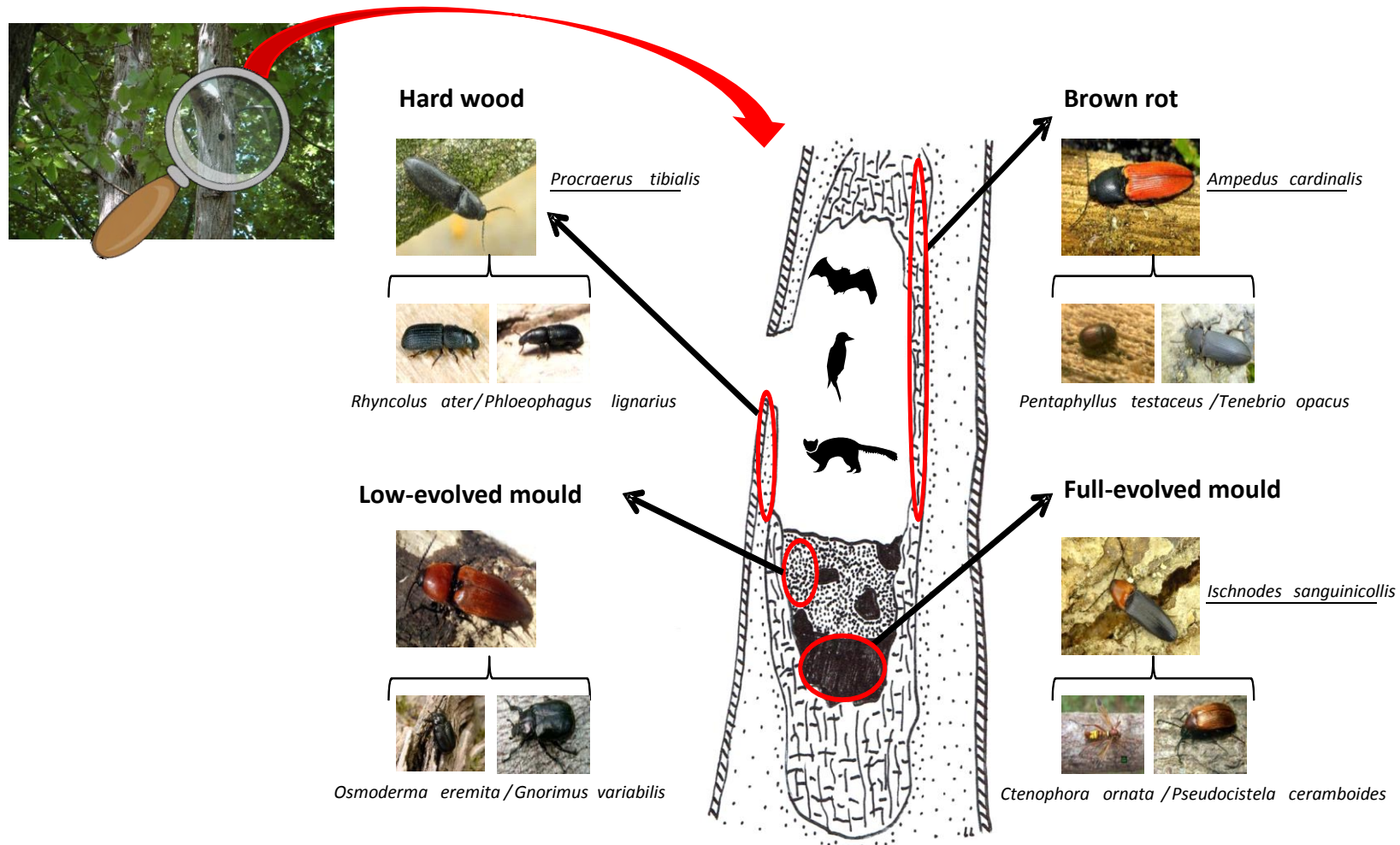
### Historical contingencies

- Legacies
- Resource gaps
- Time-lagged taxon response

## II-Relationships between TreMs and associated taxa



# A TreM is often a composite habitat and hosts several communities

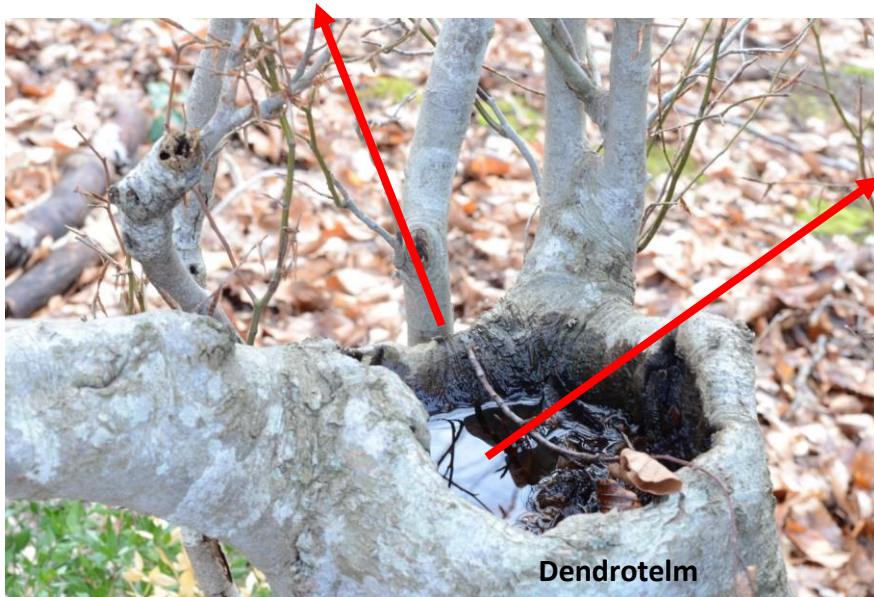


*Elateridae and their main preys; from Stokland et al. 2012 and Brustel pers. com.*

# Certain Trems host very specific species assemblages

## ☐ Mosses

- *Zygodon forsteri*
- *Anacamptodon splachnoides*



## ☐ Insects (about 15 species in Europe )

- Mainly Diptera
- Coleoptera (*Prionocyphon serricornis*)

## ☐ Fungi (Hyphomycetes)

## ☐ Flagellates, Rotifers, Nematodes

## ☐ Microcrustaceans

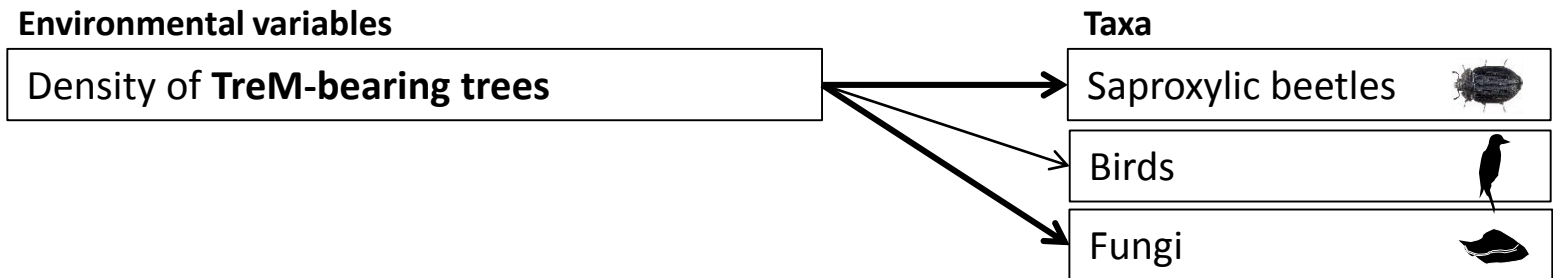
50 % of the dendrotelm-dwelling insects are strictly associated with this TreM type (Dajoz 1998)



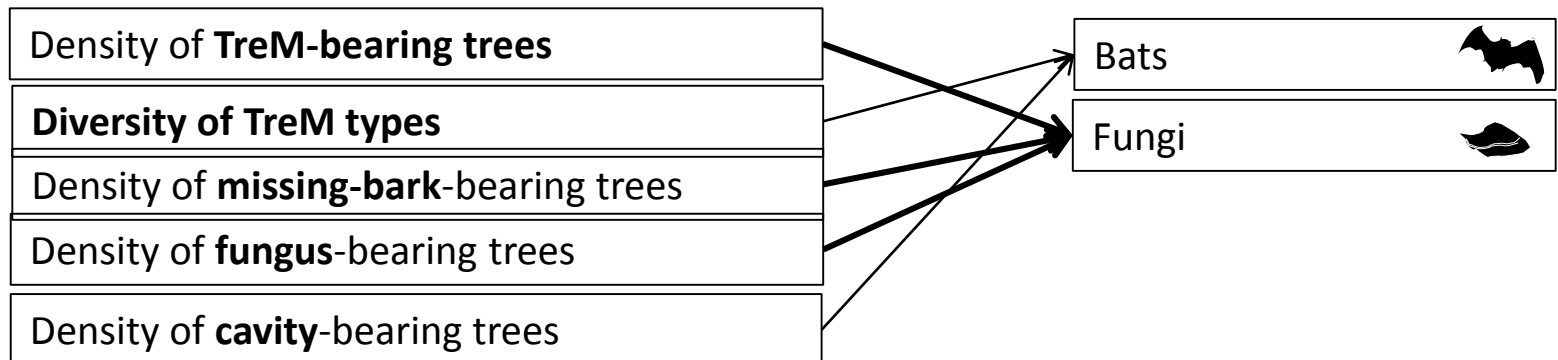
# TReM density and diversity contribute significantly to species diversity

(Larrieu et al. in prep.)

## □ Species composition (CAP)



## □ Species richness (GLMM)



Significant and positive relationships

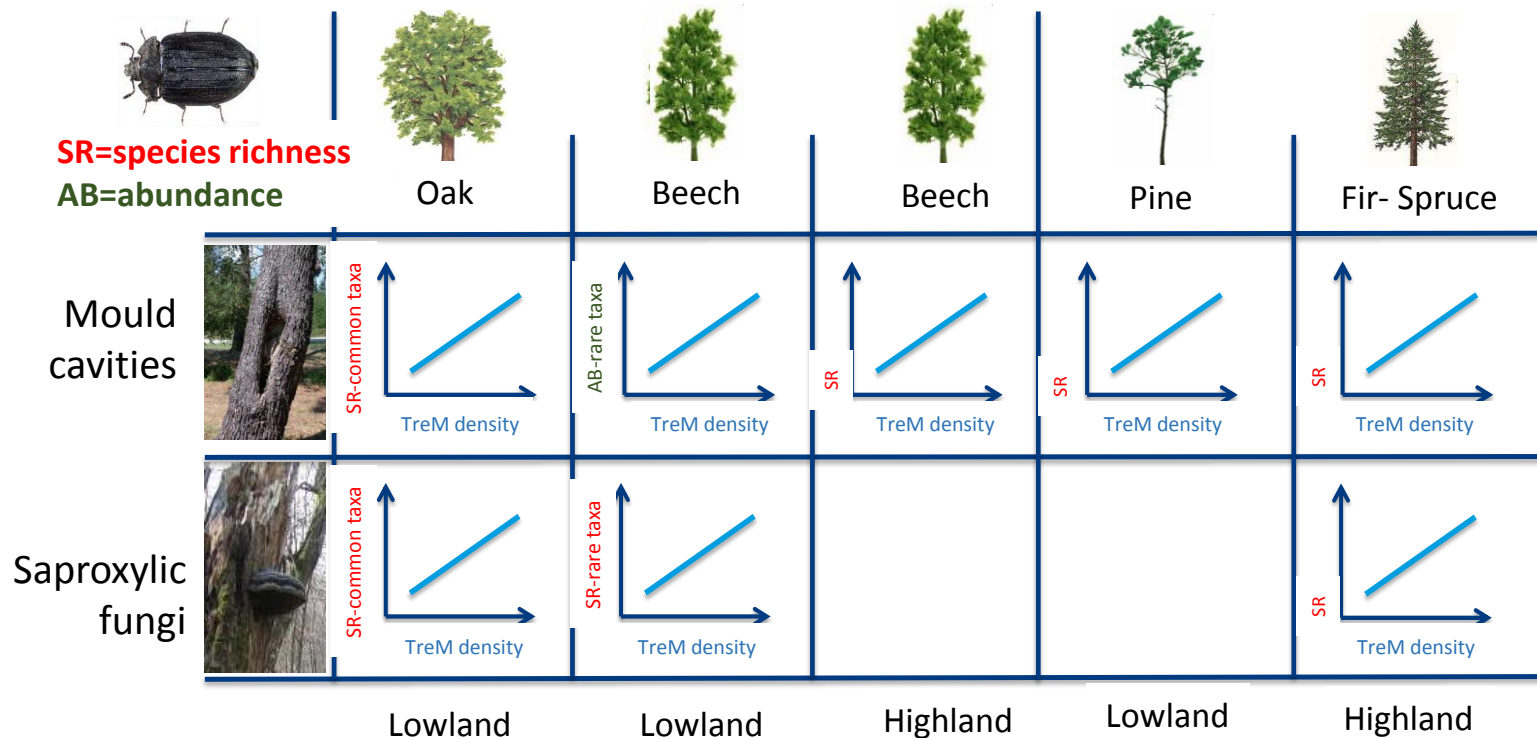
→  $p < 0.05$

→  $p < 0.001$



102 forests, harvested or not, France

# TreM-bearing tree density significantly drives saproxylic beetles diversity in many forest contexts (Bouget et al. EI 2014)



# How TReM contribute to local biodiversity depends both on forest type and taxon conservation status (Bouget et al. BC 2013)



## Contribution of TReM-bearing tree density to species richness of saproxylic beetles

### Common species

### Rare species



Oak forest

5<sup>th</sup> rank

ns



Beech forest

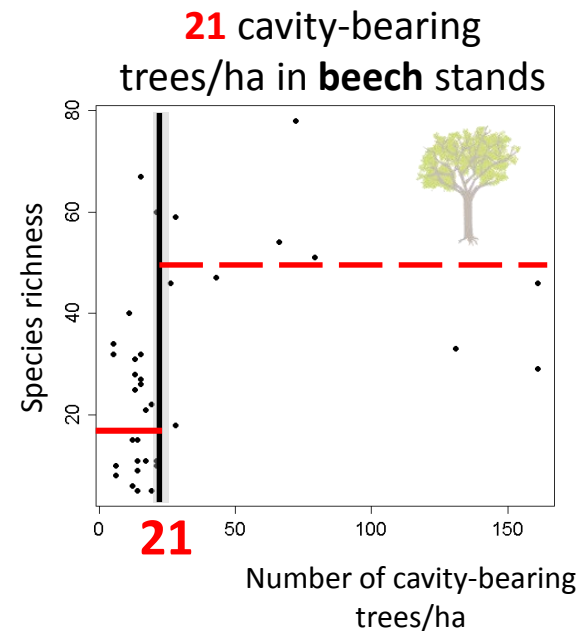
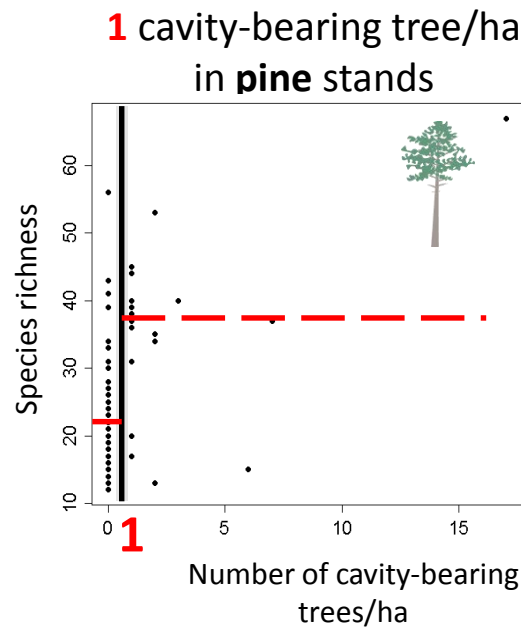
ns

1<sup>st</sup> rank

# Positive relationships between TReM density and local species richness are sometimes thresholded (Bouget et al. El 2014)



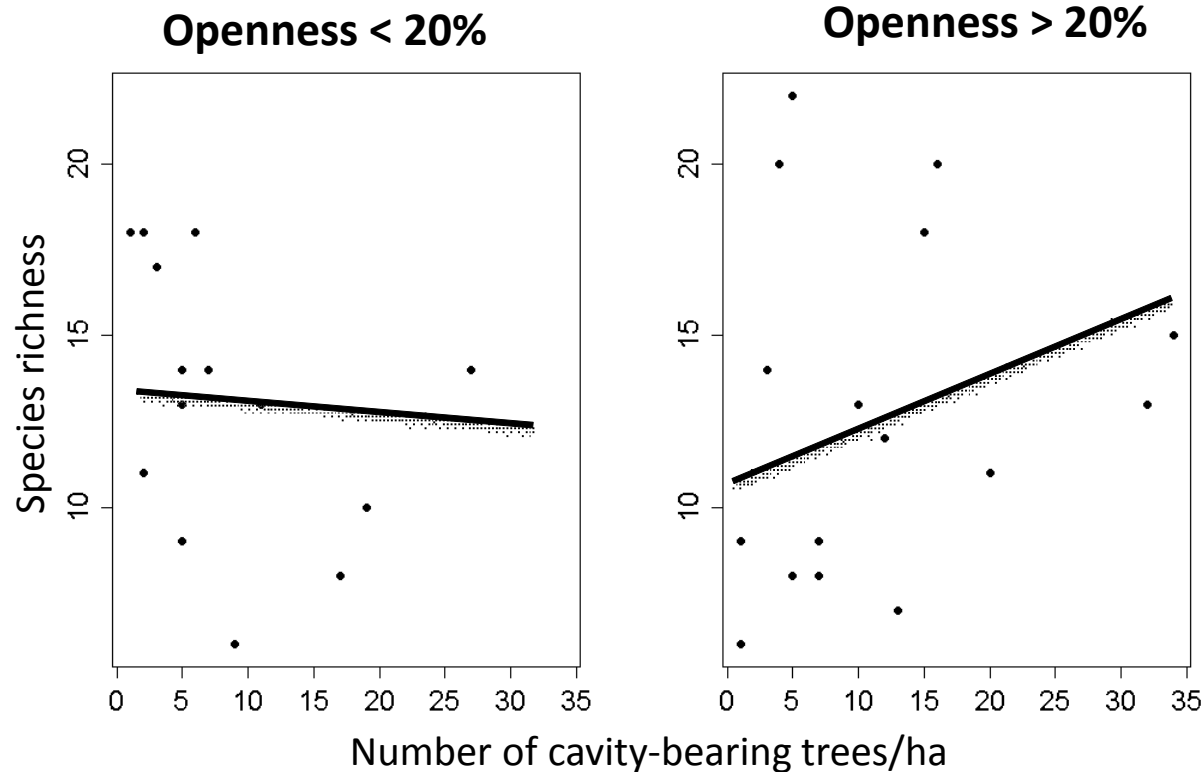
Local species richness of saproxylic beetles was, on average, higher above the thresholds





# The positive effect of increasing TReM density on saproxylic beetle diversity is affected by stand openness

(Bouget et al. EI 2014)



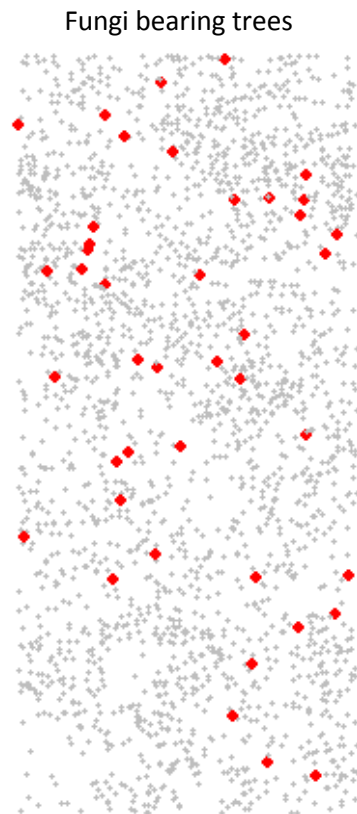
Likely effects of :

- increase of complementation resource amount (flowers,...)
- best microclimate conditions within saproxylic substrates
- beetles more active in warmer environments

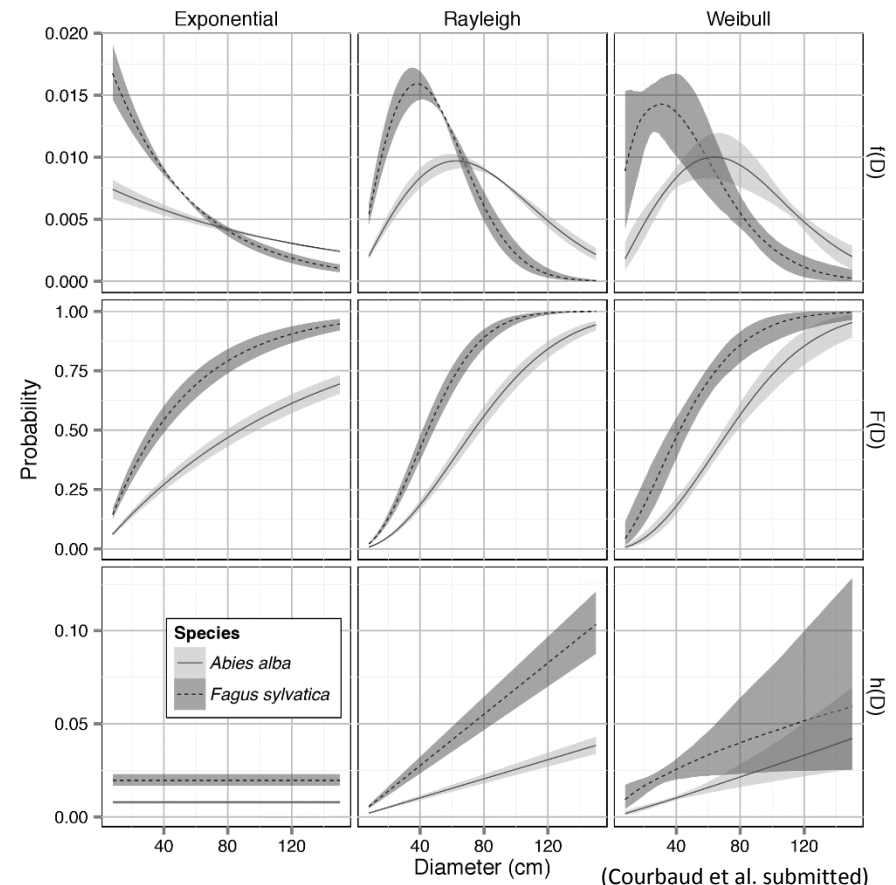


# III-Research perspectives

## Spatial distribution



## Modelling



# Ongoing research on TreM spatial patterns will take a tremendous step forward in TreM knowledge

## Main objectives

- ❑ **Mid-term: patterns of TreM spatial distribution and role on TreM-associated biodiversity**
  - In subnatural forests: TreM spatial distribution as a proxy of dispersion capacities of the TreM associated taxa
  - In managed forests: disentangle the effects of changes in both TreM density and TreM spatial distribution on TreM-associated biodiversity changes
  
- ❑ **Long-term: TreM dynamics**
  - TreM types genesis and co-occurrences (using distribution of environmental features)
  - TreM life-spans (diachronic studies)

# The modelling of TreM dynamics may help generalize results and improve practical recommendations

- ❑ **Modelling the probability of TreM formation using survival analysis methods**
- ❑ **Using tree life-history traits to understand differences of TreM dynamics among species and generalize for groups of tree species**
- ❑ **Implementing TreM dynamics in Samsara2, an individual-based model of forest dynamics**



**Prediction of TreM flows within stands, managed or not**




# IV-Related tools already available

## Typologies








## e-applications



**Fungi (and Myxomycetes)**  
Tree currently observed : 1

**Fungi (and Myxomycetes)**  
fungi (and Myxomycetes)

- 10911 - polypore   
  - 109111 - perennial    
  - 109112 - annual    
- 10912 - Myxomycete   
  - 109121 - Myxomycete    
- 10913 - Pyrenomycete   
  - 109131 - Pyrenomycete    
- 10914 - pulpy agaric   
  - 109141 - pulpy agaric    

- TreM assessment
- learning in the field
- marteloscopes







I'd like to bear  
TReMs like you...

You have to  
be patient,  
kid!

Thanks for your attention !